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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2008; month=7; day=29; hr=14; min=47; sec=35; ms=461;]

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Application No: 10527257

Version No: 5.0

Input Set:

Output Set:

Started: 2008-07-28 10:04:50.641

Finished: 2008-07-28 10:04:51.133

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 492 ms

Total Warnings: 6

Total Errors: 0

No. of SeqIDs Defined: 12

Actual SeqID Count: 12

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W 213	Artificial or Unknown found in <213> in SEQ ID (6)
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<120> TUMOR TAG AND THE USE THEREOF

<130> 186353/US

<140> 10527257

<141> 2005-03-09

<150> PCT/CN2002/000631

<151> 2002-09-09

<160> 12

<170> PatentIn version 3.4

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ctt tgc tat gac atc acc gtc atc cct aag ttc aga cct gga cca cgg	144
Leu Cys Tyr Asp Ile Thr Val Ile Pro Lys Phe Arg Pro Gly Pro Arg	
35 40 45	

tgg tgt gcg gtt caa ggc cag gtg gat gaa aag act ttt ctt cac tat	192
Trp Cys Ala Val Gln Gly Gln Val Asp Glu Lys Thr Phe Leu His Tyr	
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Asp Cys Gly Ser Lys Thr Val Thr Pro Val Ser Pro Leu Gly Lys Lys	
65 70 75 80	

cta aat gtc aca acg gcc tgg aaa gca cag aac cca gta ctg aga gag	288
Leu Asn Val Thr Thr Ala Trp Lys Ala Gln Asn Pro Val Leu Arg Glu	
85 90 95	

gtg gtg gac ata ctt aca gag caa ctg ctt gac att cag ctg gag aat	336
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tac ata ccc aag gaa ccc ctc acc ctg cag gcc agg atg tct tgt gag	384
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115 120 125	

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130 135 140	

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Gly Gln Ile Phe Leu Leu Phe Asp Ser Glu Asn Arg Met Trp Thr Thr	
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180 185 190	

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Trp Cys Ala Val Gln Gly Gln Val Asp Glu Lys Thr Phe Leu His Tyr
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Asp Cys Gly Ser Lys Thr Val Thr Pro Val Ser Pro Leu Gly Lys Lys
65 70 75 80

Leu Asn Val Thr Thr Ala Trp Lys Ala Gln Asn Pro Val Leu Arg Glu
85 90 95

Val Val Asp Ile Leu Thr Glu Gln Leu Leu Asp Ile Gln Leu Glu Asn
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Tyr Ile Pro Lys Glu Pro Leu Thr Leu Gln Ala Arg Met Ser Cys Glu
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Gly Gln Ile Phe Leu Leu Phe Asp Ser Glu Asn Arg Met Trp Thr Thr
145 150 155 160

Val His Pro Gly Ala Arg Lys Met Lys Glu Lys Trp Glu Asn Asp Lys
165 170 175

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85

90

95

Val Val Asp Ile Leu Thr Glu Gln Leu Arg Asp Ile Gln Leu Glu Asn
 100 105 110

Tyr Thr Pro Lys Glu Pro Leu Thr Leu Gln Ala Arg Met Ser Cys Glu
 115 120 125

Gln Lys Ala Glu Gly His Ser Ser Gly Ser Trp Gln Phe Ser Phe Asp
 130 135 140

Gly Gln Ile Phe Leu Leu Phe Asp Ser Glu Lys Arg Met Trp Thr Thr
 145 150 155 160

Val His Pro Gly Ala Arg Lys Met Lys Glu Lys Trp Glu Asn Asp Lys
 165 170 175

Val Val Ala Met Ser Phe His Tyr Phe Ser Met Gly Asp Cys Ile Gly
 180 185 190

Trp Leu Glu Asp Phe Leu Met Gly Met Asp Ser Thr Leu Glu Pro Ser
 195 200 205

Ala Gly Ala Pro Leu Ala Met Ser Ser Gly Thr Thr Gln Leu Arg Ala
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